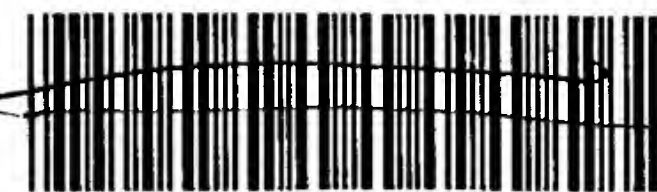


0420
 2/19/02



OIPE

RAW SEQUENCE LISTING

DATE: 02/19/2002

PATENT APPLICATION: US/10/066,960

TIME: 12:40:15

Input Set : N:\Crf3\RULE60\10066960.txt

Output Set: N:\CRF3\02192002\J066960.raw

SEQUENCE LISTING

2 (1) GENERAL INFORMATION:
 3 (i) APPLICANT: PARMA, et al.
 4 (ii) TITLE OF INVENTION: HIGH AFFINITY NUCLEIC ACID LIGANDS
 5 TO LECTINS
 6 (iii) NUMBER OF SEQUENCES: 390
 7 (iv) CORRESPONDENCE ADDRESS:
 8 (A) ADDRESSEE: Swanson & Bratschun, L.L.C.
 9 (B) STREET: 8400 E. Prentice Avenue, Suite 200
 10 (C) CITY: Englewood
 11 (D) STATE: Colorado
 12 (E) COUNTRY: USA
 13 (F) ZIP: 80111
 14 (v) COMPUTER READABLE FORM:
 15 (A) MEDIUM TYPE: Diskette, 3 1/2 diskette, 1.44 MB
 16 (B) COMPUTER: IBM pc compatible
 17 (C) OPERATING SYSTEM: MS-DOS
 18 (D) SOFTWARE: WordPerfect 6.0
 19 (vi) CURRENT APPLICATION DATA:
 C--> 20 (A) APPLICATION NUMBER: US/10/066,960
 C--> 21 (B) FILING DATE: 04-Feb-2002
 26 (C) CLASSIFICATION:
 39 (vii) PRIOR APPLICATION DATA:
 23 (A) APPLICATION NUMBER: 08/952,793
 24 (B) FILING DATE: 1999-DEC-03
 28 (A) APPLICATION NUMBER: PCT/US96/09455
 29 (B) FILING DATE: 05-JUNE-1995
 31 (A) APPLICATION NUMBER: 08/479,724
 32 (B) FILING DATE: 07-JUNE-1995
 34 (A) APPLICATION NUMBER: 08/472,256
 35 (B) FILING DATE: 07-JUNE-1995
 37 (A) APPLICATION NUMBER: 08/472,255
 38 (B) FILING DATE: 07-JUNE-1995
 40 (A) APPLICATION NUMBER: 08/477,829
 41 (B) FILING DATE: 07-JUNE-1995
 42 (viii) ATTORNEY/AGENT INFORMATION:
 43 (A) NAME: Barry J. Swanson
 44 (B) REGISTRATION NUMBER: 33,215
 45 (C) REFERENCE/DOCKET NUMBER: NEX40C/PCT
 46 (ix) TELECOMMUNICATION INFORMATION:
 47 (A) TELEPHONE: (303) 793-3333
 48 (B) TELEFAX: (303) 793-3433
 50 (2) INFORMATION FOR SEQ ID NO: 1:

ENTERED

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/066,960

DATE: 02/19/2002

TIME: 12:40:15

Input Set : N:\Crf3\RULE60\10066960.txt

Output Set: N:\CRF3\02192002\J066960.raw

```

51      (i) SEQUENCE CHARACTERISTICS:
52          (A) LENGTH: 98 base pairs
53          (B) TYPE: nucleic acid
54          (C) STRANDEDNESS: single
55          (D) TOPOLOGY: linear
W--> 56      (ii) MOLECULE TYPE: RNA
57      (ix) FEATURE:
58          (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
59      (ix) FEATURE:
60          (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
61      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
62          GGGAAAAGCG AAUCAUACAC AAGANNNNNN NNNNNNNNNN NNNNNNNNNN      50
63          NNNNNNNNNN NNNNNNNNNN NNNNGCUCCG CCAGAGACCA ACCGAGAA      98
65 (2) INFORMATION FOR SEQ ID NO: 2:
66      (i) SEQUENCE CHARACTERISTICS:
67          (A) LENGTH: 41 base pairs
68          (B) TYPE: nucleic acid
69          (C) STRANDEDNESS: single
70          (D) TOPOLOGY: linear
W--> 71      (ii) MOLECULE TYPE: RNA
72      (ix) FEATURE:
73          (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
74      (ix) FEATURE:
75          (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
76      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
77          UAAUACGACU CACUAUAGGG AAAAGCGAAU CAUACACAAG A      41
79 (2) INFORMATION FOR SEQ ID NO: 3:
80      (i) SEQUENCE CHARACTERISTICS:
81          (A) LENGTH: 24 base pairs
82          (B) TYPE: nucleic acid
83          (C) STRANDEDNESS: single
84          (D) TOPOLOGY: linear
W--> 85      (ii) MOLECULE TYPE: RNA
86      (ix) FEATURE:
87          (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
88      (ix) FEATURE:
89          (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
90      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
91          UUCUCGGUUG GUCUCUGGCG GAGC      24
93 (2) INFORMATION FOR SEQ ID NO: 4:
94      (i) SEQUENCE CHARACTERISTICS:
95          (A) LENGTH: 96 base pairs
96          (B) TYPE: nucleic acid
97          (C) STRANDEDNESS: single
98          (D) TOPOLOGY: linear
W--> 99      (ii) MOLECULE TYPE: RNA
100      (ix) FEATURE:
101          (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
102      (ix) FEATURE:

```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/066,960

DATE: 02/19/2002

TIME: 12:40:15

Input Set : N:\Crf3\RULE60\10066960.txt

Output Set: N:\CRF3\02192002\J066960.raw

```

103         (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
104     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
105         GGGAAAAGCG AAUCAUACAC AAGAAUGGUU GGCCUGGGCG CAGGCUUCGA      50
106         AGACUCGGCG GGAACGGGAA UGGCUCCGCC AGAGACCAAC CGAGAA      96
108 (2) INFORMATION FOR SEQ ID NO: 5:
109     (i) SEQUENCE CHARACTERISTICS:
110         (A) LENGTH: 98 base pairs
111         (B) TYPE: nucleic acid
112         (C) STRANDEDNESS: single
113         (D) TOPOLOGY: linear
W--> 114     (ii) MOLECULE TYPE: RNA
115     (ix) FEATURE:
116         (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
117     (ix) FEATURE:
118         (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
119     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
120         GGGAAAAGCG AAUCAUACAC AAGACAGGCA CUGAAAACUC GCGGGGAACG      50
121         AAAGUAGUGC CGACUCAGAC GCGUGCUC CGAGAGACCA ACCGAGAA      98
123 (2) INFORMATION FOR SEQ ID NO: 6:
124     (i) SEQUENCE CHARACTERISTICS:
125         (A) LENGTH: 91 base pairs
126         (B) TYPE: nucleic acid
127         (C) STRANDEDNESS: single
128         (D) TOPOLOGY: linear
W--> 129     (ii) MOLECULE TYPE: RNA
130     (ix) FEATURE:
131         (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
132     (ix) FEATURE:
133         (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
134     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
135         GGGAAAAGCG AAUCAUACAC AAGAAGUCUG GCCAAAGACU CGGCGGGAAC      50
136         GUAAAACGGC CAGAAUUGCU CCGCCAGAGA CCAACCGAGA A      91
138 (2) INFORMATION FOR SEQ ID NO: 7:
139     (i) SEQUENCE CHARACTERISTICS:
140         (A) LENGTH: 94 base pairs
141         (B) TYPE: nucleic acid
142         (C) STRANDEDNESS: single
143         (D) TOPOLOGY: linear
W--> 144     (ii) MOLECULE TYPE: RNA
145     (ix) FEATURE:
146         (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
147     (ix) FEATURE:
148         (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
149     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
150         GGGAAAAGCG AAUCAUACAC AAGAGUAGGA GGUUCCAUCA CCAGGACUCG      50
151         GCGGGAACGG AAGGUGAUGS GCUCCGCCAG AGACCAACCG AGAA      94
153 (2) INFORMATION FOR SEQ ID NO: 8:
154     (i) SEQUENCE CHARACTERISTICS:
155         (A) LENGTH: 95 base pairs

```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/066,960

DATE: 02/19/2002

TIME: 12:40:15

Input Set : N:\Crf3\RULE60\10066960.txt

Output Set: N:\CRF3\02192002\J066960.raw

```

156          (B) TYPE: nucleic acid
157          (C) STRANDEDNESS: single
158          (D) TOPOLOGY: linear
W--> 159      (ii) MOLECULE TYPE: RNA
160          (ix) FEATURE:
161              (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
162          (ix) FEATURE:
163              (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
164          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
165      GGGAAAAGCG AAUCAUACAC AAGAACAAGG AUCGAUGGCG AGCCGGGGAG      50
166      GGCUCGGCGG GAACGAAAUUC UGCUCCGCCA GAGACCAACC GAGAA      95
168 (2) INFORMATION FOR SEQ ID NO: 9:
169      (i) SEQUENCE CHARACTERISTICS:
170          (A) LENGTH: 97 base pairs
171          (B) TYPE: nucleic acid
172          (C) STRANDEDNESS: single
173          (D) TOPOLOGY: linear
W--> 174      (ii) MOLECULE TYPE: RNA
175          (ix) FEATURE:
176              (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
177          (ix) FEATURE:
178              (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
179          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
180      GGGAAAAGCG AAUCAUACAC AAGAUUGGGC AGGCAGAGCG AGACCGGGGG      50
181      CUCGGCGGGA ACGGAACAGG AAUGCUCGCG CAGAGACCAA CCGAGAA      97
183 (2) INFORMATION FOR SEQ ID NO: 10:
184      (i) SEQUENCE CHARACTERISTICS:
185          (A) LENGTH: 97 base pairs
186          (B) TYPE: nucleic acid
187          (C) STRANDEDNESS: single
188          (D) TOPOLOGY: linear
W--> 189      (ii) MOLECULE TYPE: RNA
190          (ix) FEATURE:
191              (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
192          (ix) FEATURE:
193              (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
194          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
195      GGGAAAAGCG AAUCAUACAC AAGAAAGGGA UGGGAUUGGG ACGAGCGGCC      50
196      AAGACUCGGC GGGAACGAAG GGUGCUCGCG CAGAGACCAA CCGAGAA      97
198 (2) INFORMATION FOR SEQ ID NO: 11:
199      (i) SEQUENCE CHARACTERISTICS:
200          (A) LENGTH: 96 base pairs
201          (B) TYPE: nucleic acid
202          (C) STRANDEDNESS: single
203          (D) TOPOLOGY: linear
W--> 204      (ii) MOLECULE TYPE: RNA
205          (ix) FEATURE:
206              (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
207          (ix) FEATURE:

```

RAW SEQUENCE LISTING

DATE: 02/19/2002

PATENT APPLICATION: US/10/066,960

TIME: 12:40:15

Input Set : N:\Crf3\RULE60\10066960.txt

Output Set: N:\CRF3\02192002\J066960.raw

```

208         (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
209     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
210         GGGAAAAGCG AAUCAUACAC AAGACUCGGC GGGAACGAAA GUGUCAUGGU       50
211         AGCAAGUCCA AUGGUGGACU CUGCUCGCC AGAGACCAAC CGAGAA           96
213 (2) INFORMATION FOR SEQ ID NO: 12:
214     (i) SEQUENCE CHARACTERISTICS:
215         (A) LENGTH: 98 base pairs
216         (B) TYPE: nucleic acid
217         (C) STRANDEDNESS: single
218         (D) TOPOLOGY: linear
W--> 219     (ii) MOLECULE TYPE: RNA
220     (ix) FEATURE:
221         (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
222     (ix) FEATURE:
223         (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
224     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
225         GGGAAAAGCG AAUCAUACAC AAGACUCGGC GGGAACGUGA AGUGGGUAGG       50
226         UAGCUGAAGA CGGUCUGGGC GCCAGCUCCG CCAGAGACCA ACCGAGAA       98
228 (2) INFORMATION FOR SEQ ID NO: 13:
229     (i) SEQUENCE CHARACTERISTICS:
230         (A) LENGTH: 99 base pairs
231         (B) TYPE: nucleic acid
232         (C) STRANDEDNESS: single
233         (D) TOPOLOGY: linear
W--> 234     (ii) MOLECULE TYPE: RNA
235     (ix) FEATURE:
236         (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
237     (ix) FEATURE:
238         (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
239     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
240         GGGAAAAGCG AAUCAUACAC AAGAAAGGGA UGGGAUUGGG ACGAGCGGCC       50
241         AAGACUCGGC GGGAACGAAG GUUCCGCUCC GCCAGAGACC AACCGAGAA       99
243 (2) INFORMATION FOR SEQ ID NO: 14:
244     (i) SEQUENCE CHARACTERISTICS:
245         (A) LENGTH: 98 base pairs
246         (B) TYPE: nucleic acid
247         (C) STRANDEDNESS: single
248         (D) TOPOLOGY: linear
W--> 249     (ii) MOLECULE TYPE: RNA
250     (ix) FEATURE:
251         (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
252     (ix) FEATURE:
253         (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
254     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
255         GGGAAAAGCG AAUCAUACAC AAGACUCGGC GGGAACGAAG UGUGUGAGUA       50
256         ACGAUCACUU GGUACUAAAA GCCCGCUCCG CCAGAGACCA ACCGAGAA       98
258 (2) INFORMATION FOR SEQ ID NO: 15:
259     (i) SEQUENCE CHARACTERISTICS:
260         (A) LENGTH: 100 base pairs

```


VERIFICATION SUMMARY

DATE: 02/19/2002

PATENT APPLICATION: US/10/066,960

TIME: 12:40:16

Input Set : N:\Crf3\RULE60\10066960.txt

Output Set: N:\CRF3\02192002\J066960.raw

L:20 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:21 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:56 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:56 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=1
L:71 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:71 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=2
L:85 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:85 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=3
L:99 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:99 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=4
L:114 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:114 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=5
L:129 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:129 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=6
L:144 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:144 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=7
L:159 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:159 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=8
L:174 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:174 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=9
L:189 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:189 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=10
L:204 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:204 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=11
L:219 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:219 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=12
L:234 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:234 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=13
L:249 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:249 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=14
L:264 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:264 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=15
L:279 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:279 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=16
L:295 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:295 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=17
L:310 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:310 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=18
L:325 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:325 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=19
L:340 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:340 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=20
L:355 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:355 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=21
L:370 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:370 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=22
L:385 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:385 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=23

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/066,960

DATE: 02/19/2002

TIME: 12:40:16

Input Set : N:\Crf3\RULE60\10066960.txt

Output Set: N:\CRF3\02192002\J066960.raw

L:400 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:400 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=24
L:415 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:415 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=25
L:430 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:430 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=26
L:445 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:445 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=27
L:460 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:460 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=28
L:475 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:475 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=29
L:490 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:490 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=30
L:505 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:505 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=31
L:520 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:520 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=32
L:535 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:535 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=33
L:550 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:550 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=34
L:565 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:565 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=35
L:580 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:580 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=36
L:595 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:595 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=37
L:610 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:610 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=38
L:625 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:625 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=39
L:640 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:640 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=40
L:655 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:655 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=41
L:670 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:670 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=42
L:685 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:685 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=43
L:700 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:700 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=44
L:715 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:715 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=45
L:730 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:730 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=46
L:745 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:745 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=47
L:760 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/066,960

DATE: 02/19/2002

TIME: 12:40:16

Input Set : N:\Crf3\RULE60\10066960.txt

Output Set: N:\CRF3\02192002\J066960.raw

L:760 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=48
L:775 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=49
L:790 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=50